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piece 1, NC_000913, ydiP_ydiQ+, config: linear, direction: +, begin: 1777296, end: 1777660

5' *1777300 * *1777310 * *1777320 * *1777330 * *1777340 * *1777350 * *1777360 * *1777370 * 3'
- thr - gly - val - ile - lys - thr - ala - leu - ile - his - asp - asn - leu - pro - ile - pro -
- leu - ala - leu - ser - lys - gln - arg - fMet - thr - thr - ser - leu - phe - his - glu - gln - ala - lys -
- trp - arg - tyr - gln - asn - ser - val - asp - thr - fMet - ser - lys - gln - lys - gln - tyr - met - pro -
asp - lys - ile -

... NC_000913.ydiP p10 1.0 bits [###> orf 17 codons



{ sd-(10)-ir 1777324 Gap 2.7 bits
sd-ir 1777324 ydiP_ydiQ+ total 5.1 bits

p35 5.6 bits

{ p35-(22)-p10 1777340 Gap 2.3 bits
p35-p10 1777340 total 4.2 bits

5' *1777380 * *1777390 * *1777400 * *1777410 * *1777420 * *1777430 * *1777440 * *1777450 * 3'
- fMet - tyr - cys - gly - tyr - thr - ala - phe - ser - glu - ser - fMet - tyr - gly -
- arg - arg - val - leu - arg - leu - tyr - asn - arg - val - fMet - gln - asp - cys - thr - val -

p35 2.3 bits

sd 5.8 bits { ... ir

p10 5.5 bits

{ p35-(26)-p10 1777434 Gap 3.7 bits p10 2.5 bits

|-----| p35-p10 1777434 total 4.4 bits

{ p35-(21)-p10 1777453 Gap 3.3 bits
p35-p10 1777453 total 4.7 bits

5' *1777460 * *1777470 * *1777480 * *1777490 * *1777500 * *1777510 * *1777520 * *1777530 * 3'
- his - gly - ile - thr - ile - leu - pro - gln - ile - ala - leu - leu - phe - fMet - his - phe - cys - phe - ser -
- met - glu - fMet - glu - ile - thr - ile - leu - pro - gln - fMet - his - phe - cys - phe - ser - asp - arg - ile - phe -
- ala - leu - leu - ala - arg - leu - ala - arg - lys - thr - fMet - gln - asp - cys - trp - gln - gly - lys - gln -

... --| sd-(12)-ir 1777459 Gap 4.0 bits
... --| sd-ir 1777459 ydiP_ydiQ+ total 6.2 bits

[###> orf 3 codons

ir ydiP_ydiQ+ p35 4.7 bits

p10 2.3 bits

{ p35-(23)-p10 1777507 Gap 1.4 bits
p35-p10 1777507 total 5.5 bits

p35 3.8 bits

p10 2.8 bits

{ p35-(23)-p10 1777518 Gap 1.4 bits
p35-p10 1777518 total 5.1 bits

5' *1777540 * *1777550 * *1777560 * *1777570 * *1777580 * *1777590 * *1777600 * *1777610 * 3'
- fMet - leu - arg - arg - lys - pro - arg - thr - ala - ile - ala - his - tyr - phe - gln - ala - ile - phe - thr - phe - his - arg - arg - trp - phe - arg -
- ala - cys - ser - val - glu - asn - pro - ala - pro - leu - ser - his - thr - ile - phe - arg - pro - phe - leu - pro - ser - ile - gly - asp - gly - ser - val -
- leu - ala - pro - ser - lys - thr - pro - his - arg - tyr - arg - thr - ile - phe - ser - gly - his - phe - tyr - leu - pro - ser - glu - met - val - pro - tyr -

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The diagram illustrates a bacterial promoter region with the following details:

- Start Sites:** Five promoters are indicated by asterisks (*): *1777620, *1777630, *1777640, *1777650, and *1777660.
- Gene Products:** The genes are labeled as follows:
 - *1777620: a t g c g a c t c a c a g g a g a a t c a t c g a a a t a a t a a c c t g c t t 3'
 - *1777630: met - arg - leu - thr - gly - glu - ile - met - lys - ile - ile - thr - cys -
 - *1777640: cys - asp - ser - gln - glu - lys - ser -
 - *1777650: ala - thr - his - arg - arg - asn - his - glu - asn - asn - asn - leu - leu -
 - *1777660: (no sequence shown)
- Promoter Elements:**
 - A green box labeled "sd" represents the -35/-10 promoter elements.
 - A red box labeled "ir ydiP_ydiQ+" represents the inverted repeat element.
 - A blue box labeled "sd-(8)-ir 1777641 Gap 2.4 bits" indicates a gap between the -35/-10 element and the inverted repeat.
 - A blue box labeled "sd-ir 1777641 ydiP_ydiQ+ total 17.4 bits" indicates the total length of the -35/-10 element and the inverted repeat.
- Transcription Direction:** Arrows indicate transcription from left to right for all genes.
- NC_000913.ydiQ:** A label at the end of the sequence indicates the reference genome and gene name.